

FIG. 1A

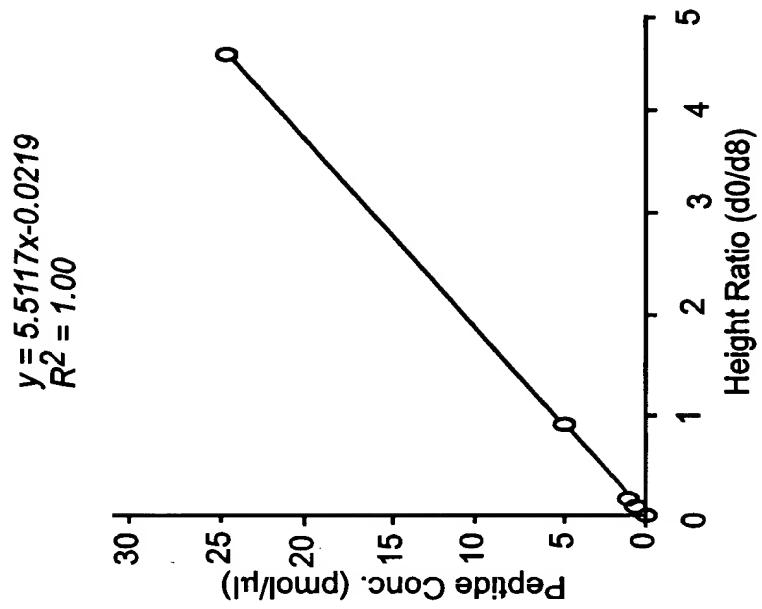
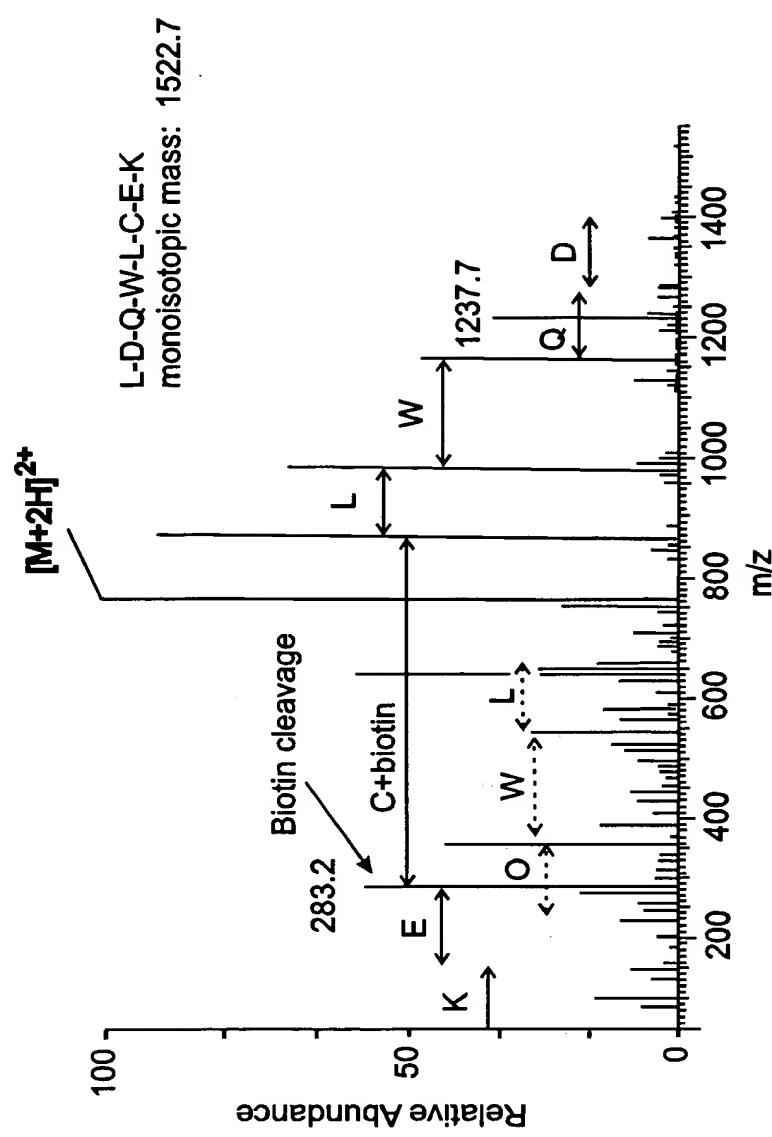


FIG. 1B

FIG. 2



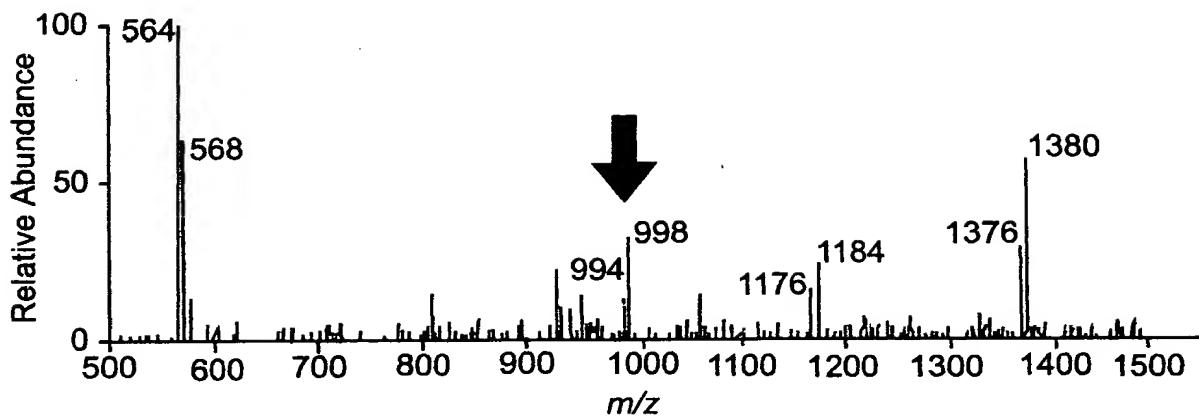


FIG. 3A

FIG. 3B

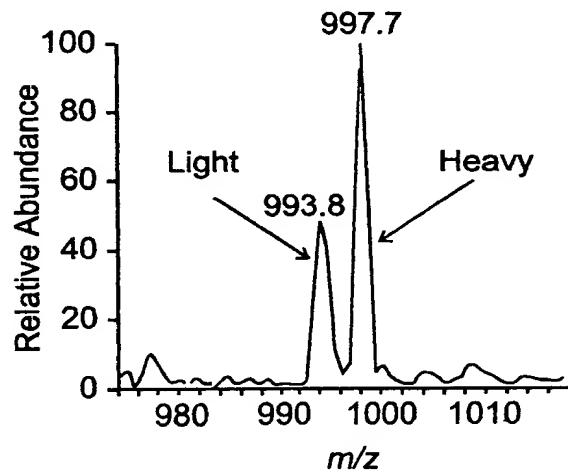
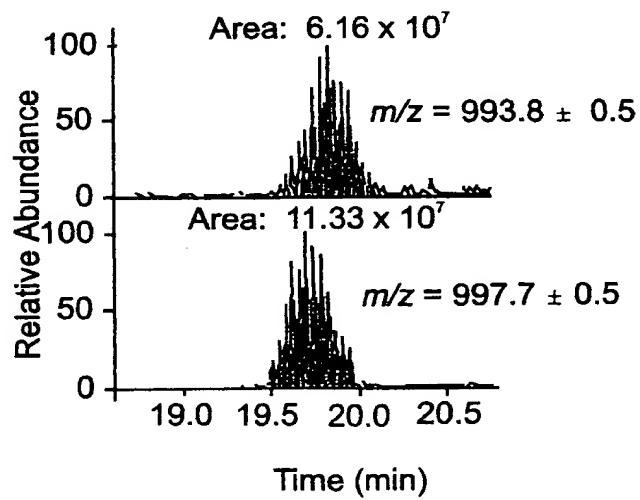


FIG. 3C



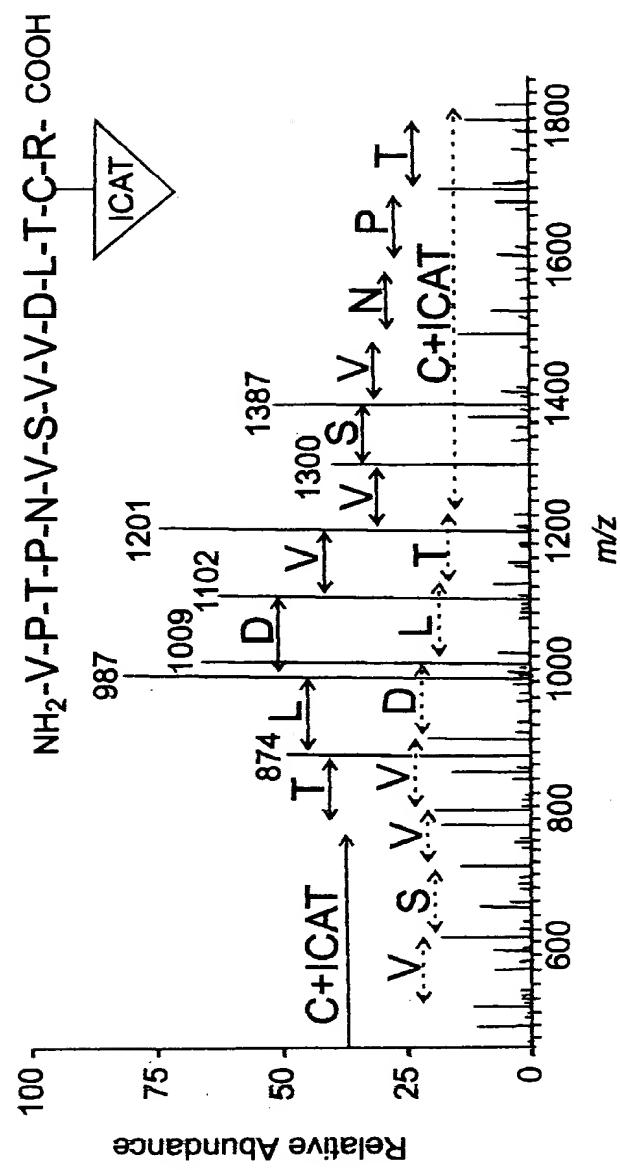


FIG. 4A

#	<u>Rank/Sp</u>	<u>(M+H)+C*10^4</u>	<u>Ions</u>	<u>Reference</u>	<u>Peptide</u>
1.	1/ 1	1994.3 4.4675	17/26	G3P_RABIT	(R)VPTPNVSVVVDLTC#R (SEQ ID NO:60)
2.	2/ 403	1995.1 2.7366	13/34	SLTRNGL	(E)LGKPVLTANQVTIWEGLR (SEQ ID No:61)
3.	3/ 3	1995.0 2.6591	16/36	FLP_LACCA	(N)LANPNVYVTETLTAATVCTI (SEQ ID NO:62)
4.	4/ 209	1995.0 2.6335	14/36	A42912	(Y)ALLPSPDAEGPHGQFVTDK (SEQ ID NO:63)
5.	5/ 381	1995.1 2.4634	13/38	H69373	(L)ALLVLVAPAMAAGNGEDLRLN (SEQ ID NO:64)

## FIG. 4B

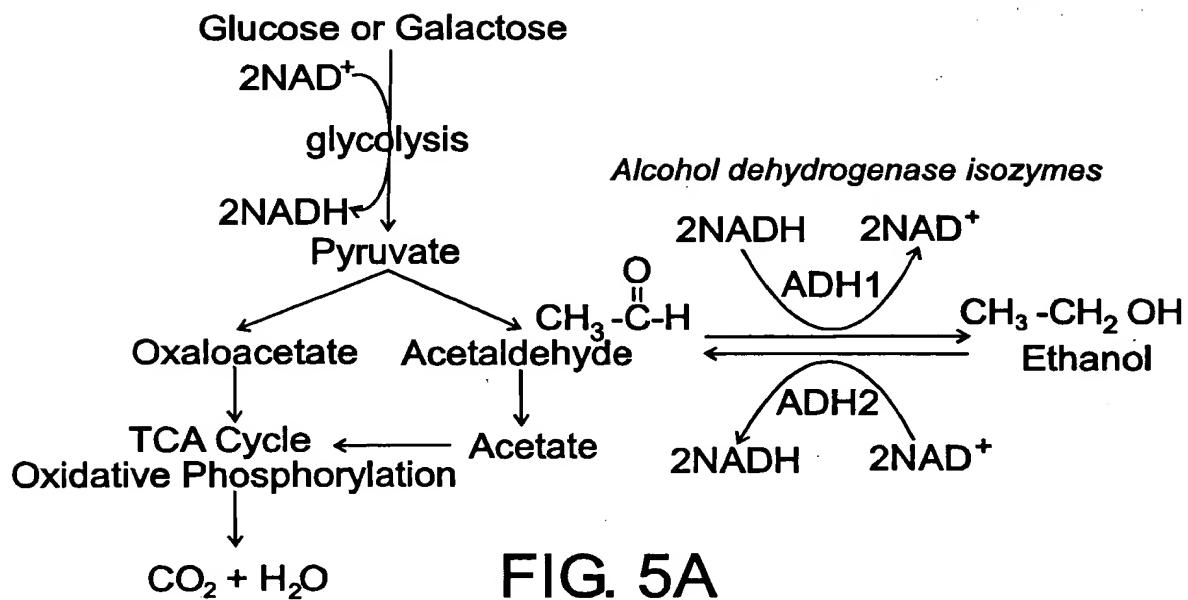


FIG. 5A

ADH1 : YSVC#HTDLHAWHGDWPLPVK

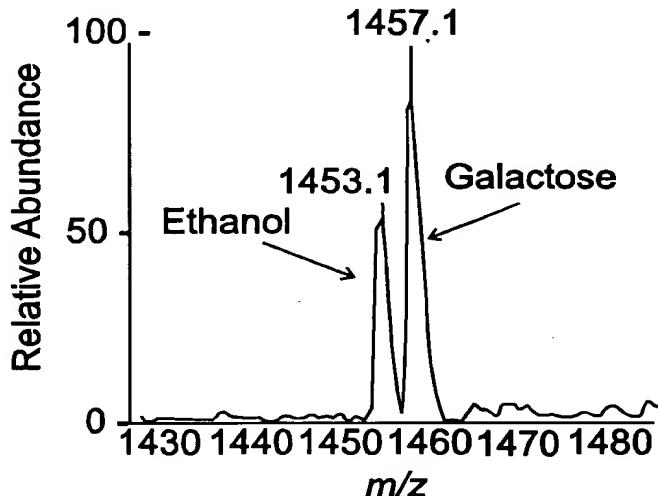


FIG. 5B

ADH2 : YSVC#HTDLHAWHGDWPLPTK

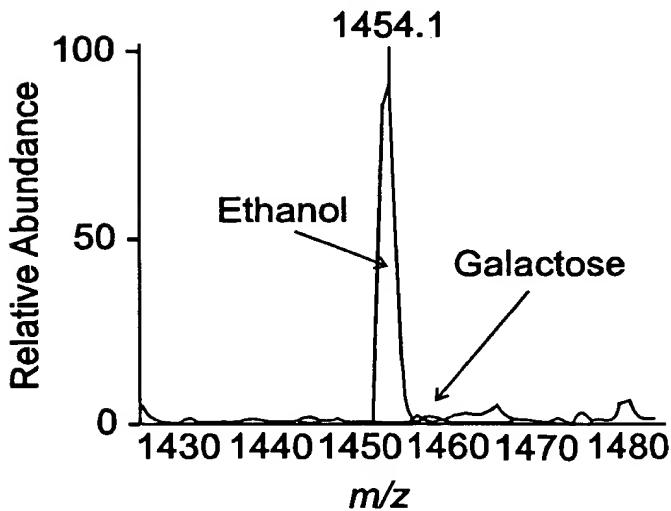


FIG. 5C

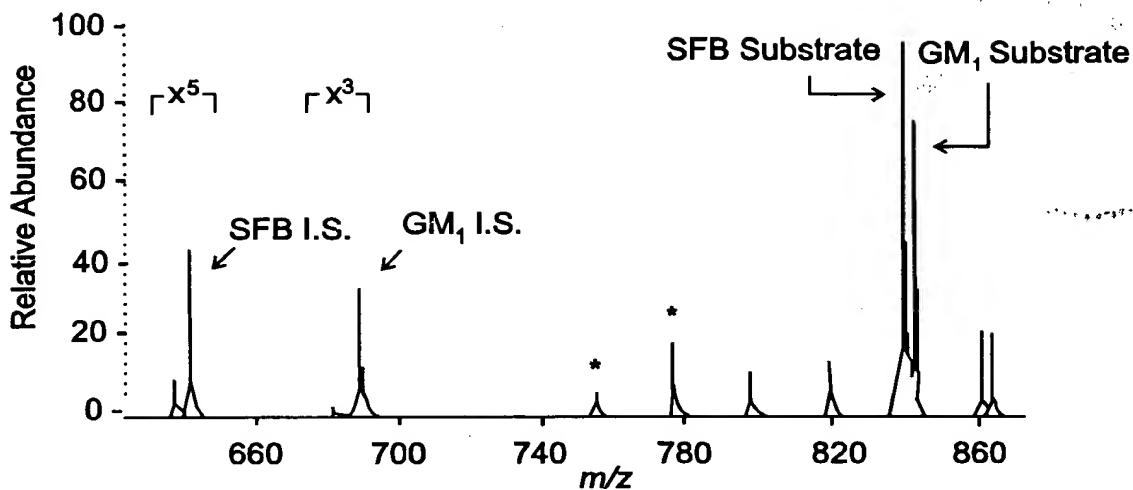


FIG. 6A

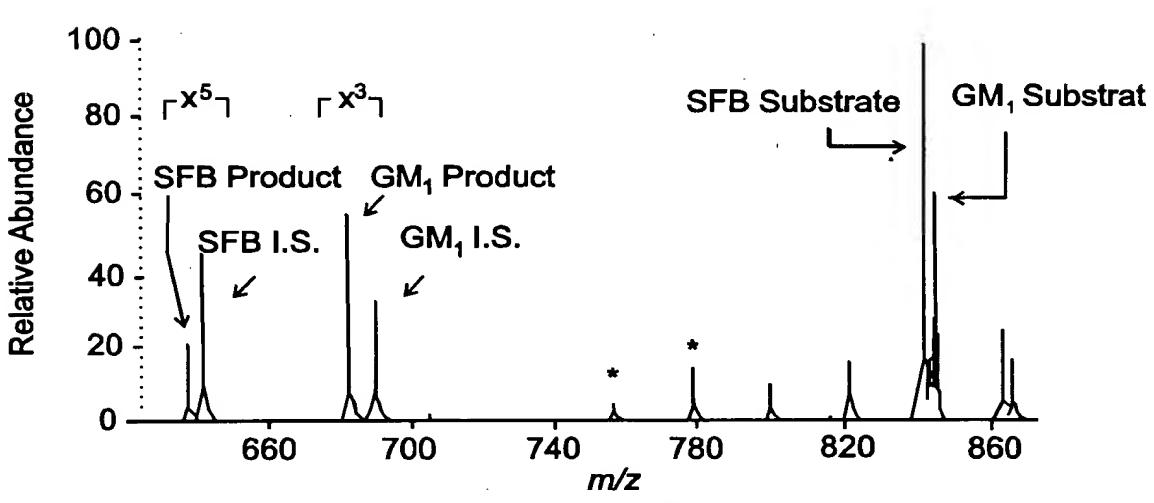


FIG. 6B

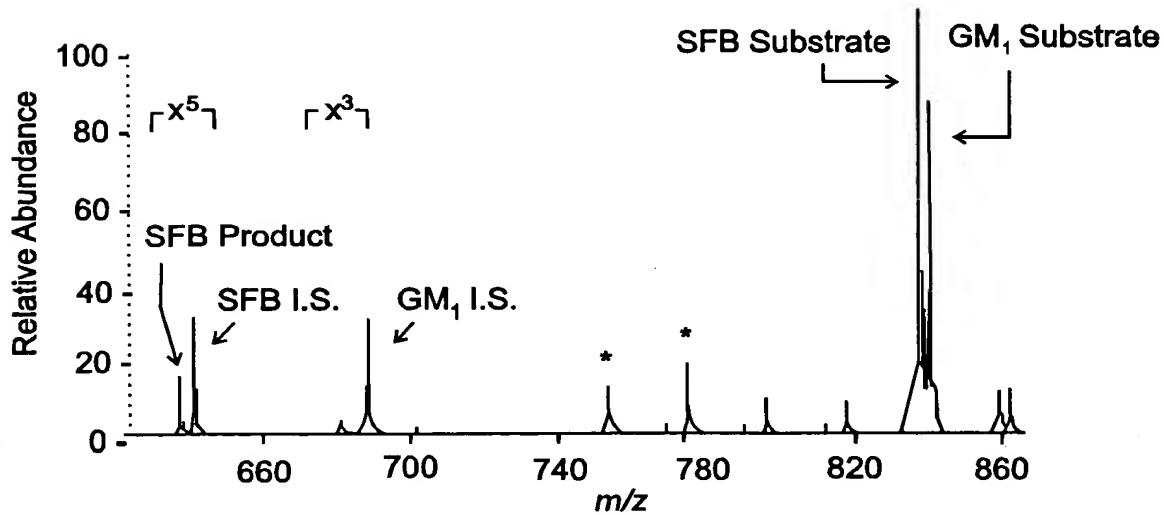
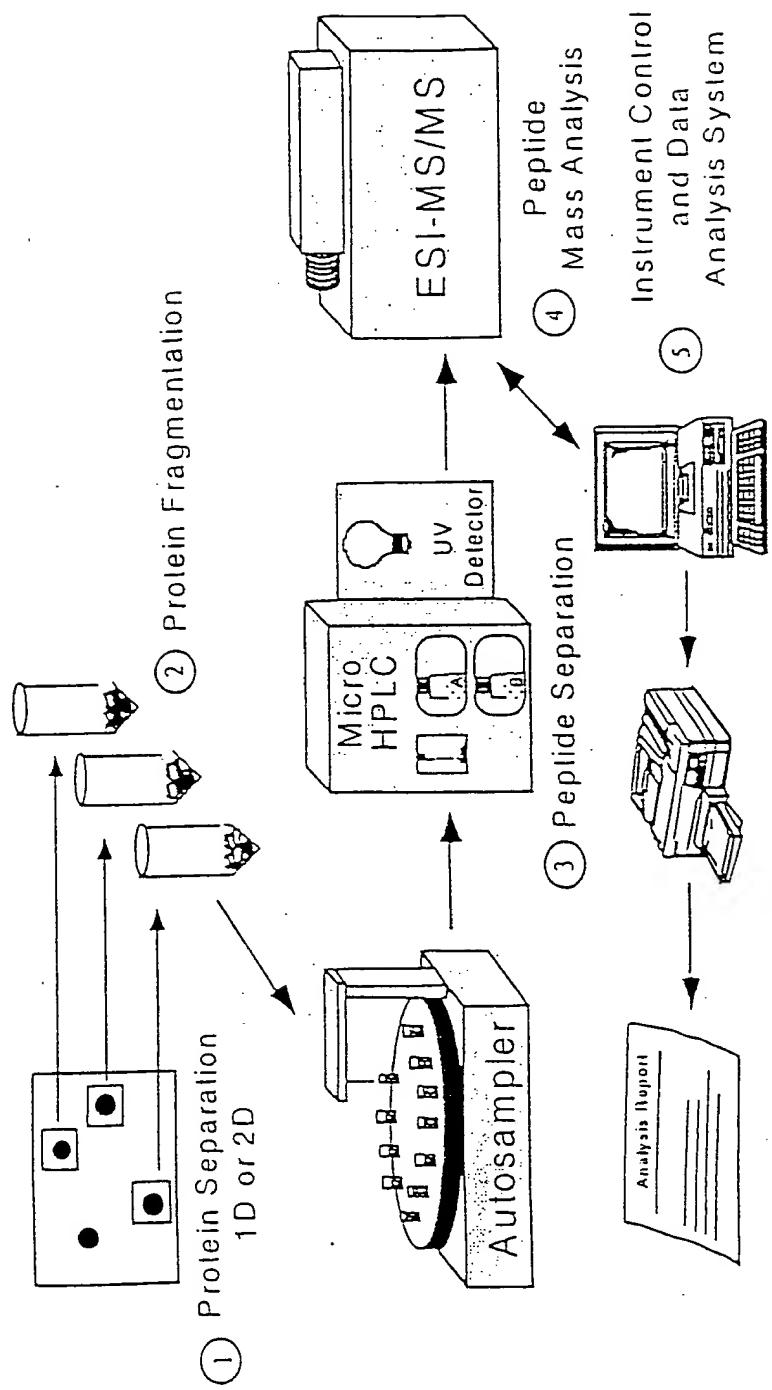


FIG. 6C

FIG. 7



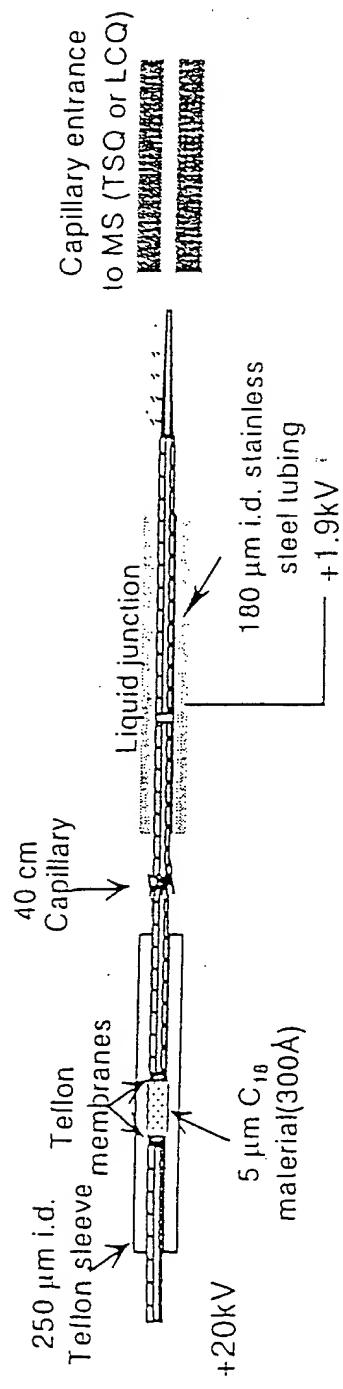


FIG. 8A

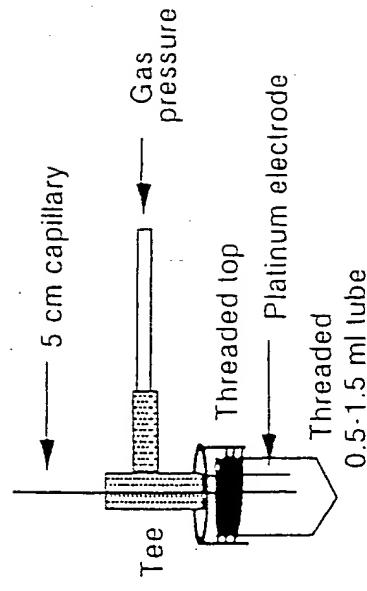


FIG. 8B